```
gtccaaggtc cagaaagcaa aaagggtgcc ccgtttctcc tgggaaggct tccccgtcag    2460

tgatttctgt aaccggaccc tgccctgaca cagcgtcatt ggactaccca gcagacagta    2520

gactccactc taaacccgct tcttgcggtc agttgctgtc cttcagtgtg tgtaagcagt    2580

ggccagacag caccctggg tgtcatttca agactctctc accttggtct gctttacgtt    2640

tggtttgatt tggtttgttc tggtttttga dacgaggcct ttcactggaa cctggcactc    2700

agtatttaga ctgcccagcc agctagcctc agagaatgca tctgcgtatg cttgcctggc    2760

gctggaattc ggtgcacatg gctttgatgt gtaccgggga tcagacacag atgtttcatg    2820

agtgcagtgc atgcctgtta gtggtagagc tc                                  2852
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<210>   8
<211>   41
<212>   PRT
<213>   Homo sapiens

<400>   8

```
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1               5                   10                  15


Pro Thr Leu Leu Leu Leu Pro Val Arg Gly Pro Arg Pro Arg Ser Phe
            20                  25                  30


Ser Ala Pro Phe Ser Ser Gln Asp Ser
        35                  40
```

<210>   9
<211>   20
<212>   DNA
<213>   Artificial

<220>
<223>   SYNTHETIC PRIMER

<220>
<221>   misc_feature
<222>   (1)..(20)
<223>   SYNTHETIC PRIMER

<400>   9
ttcacaccac tctgtcgaac   20

<210>   10
<211>   20
```

```
<220>
<221>  misc_feature
<222>  (1)..(19)
<223>  construct

<400>  36                                                    19
aggcataaaa aagtcggtc


<210>  37
<211>  15
<212>  PRT
<213>  Unknown

<220>
<223>  conserved sequence


<220>
<221>  DOMAIN
<222>  (1)..(15)
<223>  conserved sequence

<400>  37

Pro Arg Pro Arg Ser Phe Ser Ala Pro Phe Ser Ser Gln Asp Ser
1                5                   10                  15
```